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REPLACEMENT SHEET
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Fig 1A

S-GACCCTAGAT CCCAAGATCT CCAAGGATTT GGTGGCATAC CCACTCCAGC ACACAGAAGC <u>ATG</u>AGGTTCA start codon 5'-UTS →

TGACTCTCCT CTTCCTGACA GCTCTGGCAG GAGCCCTGGT CTGTGCCT 1 AT GATCCAGAGG CCGCCTCTGC exon 2 intron 1 exon 1

CCCAGGATCG GGGAACC II CTT GCCATGAAGC ATCAGCAGCT CAAAAGGAAA ATGCAGGTGA AGACCCAGGG exon 3 intron 2

TTAGCCAGAC AGGCACCAAA GCCAAGGAAG CAGAGATCCA GCCTTCTGG 1 A AAAAGGCCTA GACGGAGCAA exon 4 intron 3

AAAAAGCTGT GGGGGACTC GGAAAACTAG GAAAAGATGC AGTCGAAGAT CTAGAAAGCG TGGGTAAAGG

AGCCGTCCAT GACGTTAAAG ACGTCCTTGA CTCAGTACTA TAGCTGTAAG GAGAAGCTGA GAAA1GATAC stop codon exon 5

← 3'-UTS CCAGGAGCAG CAGGCTTTAC GTTTTCAGCC TAAAACCT

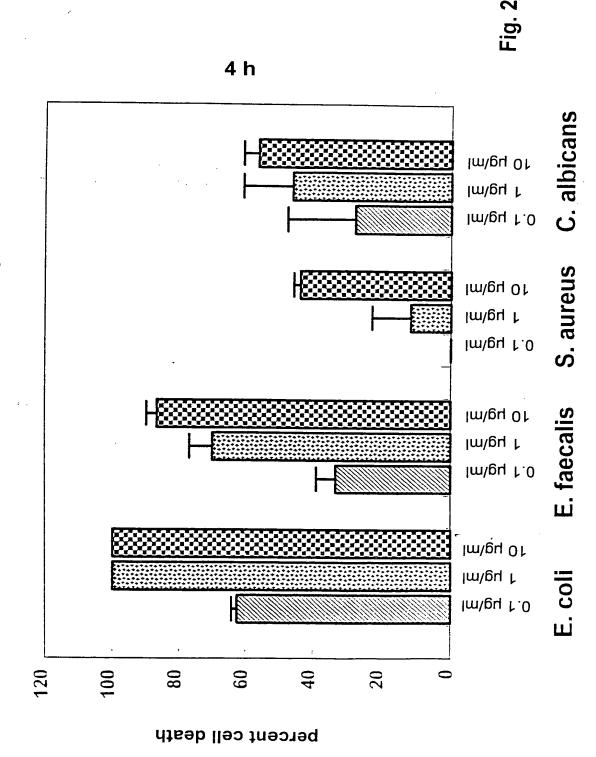
Fig 1B

QRSSLLEKGL MRFMTLLFLT ALAGALVCAY DPEAASAPGS GNPCHEASAA QKENAGEDPG LARQAPKPRK DGAKKAVGGL GKLGKDAVED LESVGKGAVH DVKDVLDSVL

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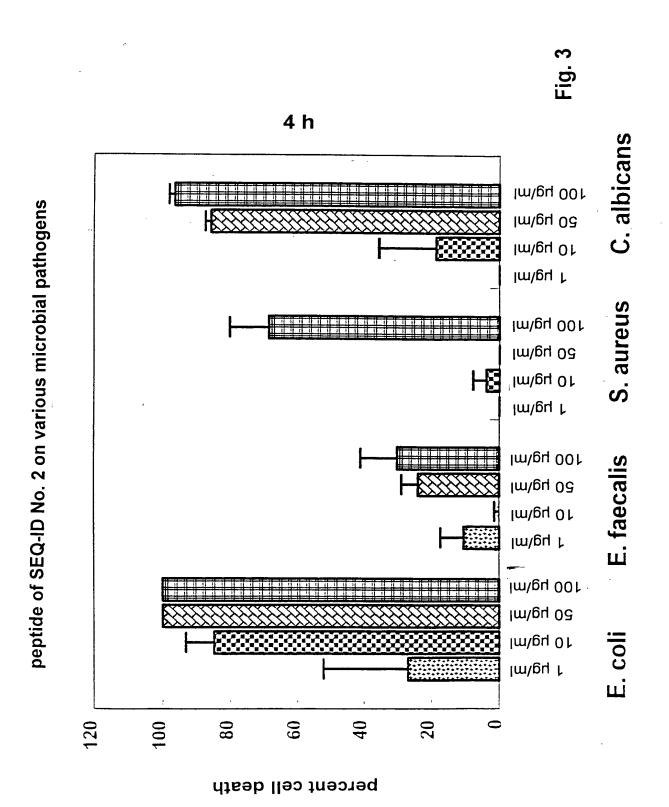


DCD-eGFP on various microbial pathogens

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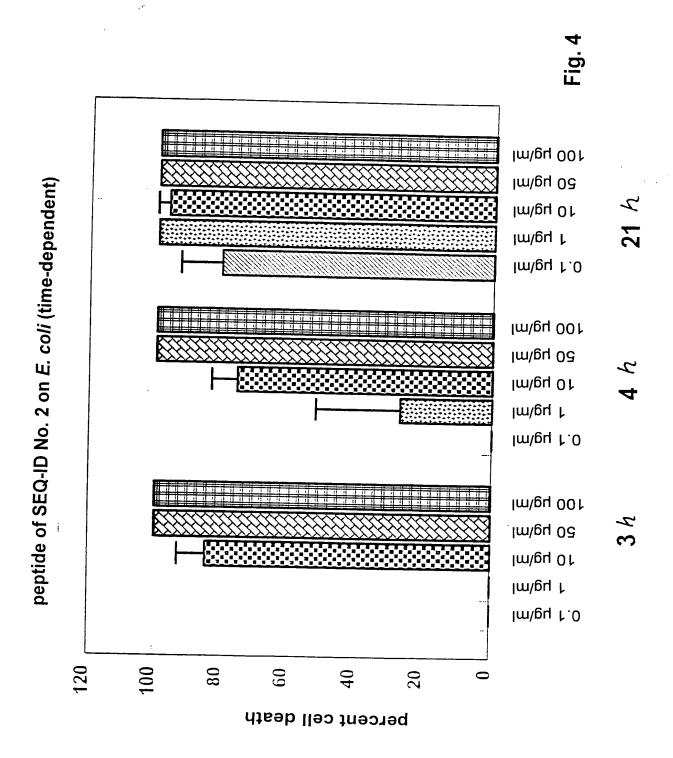
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